Structure and dynamics of single-isoform recombinant neuronal human tubulin

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SUPPLEMENTAL TABLE 1. Refinement statistics and model geometry

Refinement statistics and model geometry from REFMAC v5.8 (29) and MolProbity (51).

Resolution for refinement	4 Å
Map sharpening B factor	-180
FSC _{average} ^a	0.79
Rms deviations (bonds)(Å ²)	0.008
Rms deviations (angles)(Å ²)	1.438
Molprobity Score	1.75 (100 th percentile)
Clashscore, all atoms	7.39 (100th percentile)
Poor rotamers (%)	1.18%
Ramachandran plot favoured (%)	95.75%
Ramachandran outliers (%)	0.12%

^aaFSCaverage = $\Sigma(N_{shell} FSC_{shell})/\Sigma(N_{shell})$, where FSC_{shell} is the FSC in a given shell, N_{shell} is the number of structural factors in the shell. FSC_{shell} = $\Sigma(F_{model} F_{EM})/(\sqrt{\Sigma(|F|2_{model})})\sqrt{\Sigma(F^2_{EM})}$

SUPPLEMENTAL MOVIE 1

Dynamics of recombinant human α1A/βIII microtubules imaged by darkfield microscopy.

Dynamic recombinant human $\alpha 1A/\beta III$ microtubules at 9 μM tubulin imaged by darkfield microscopy. Blue marks the GMPCPP seed. Recorded at 12 frames per minute, played back at 30 frames per second.

SUPPLEMENTAL MOVIE 2

Dynamics of heterogeneous brain microtubules imaged by darkfield microscopy.

Dynamic brain microtubules at 9 μ M tubulin imaged by darkfield microscopy. Blue marks the GMPCPP seed. Recorded at 12 frames per minute, played back at 30 frames per second.

SUPPLEMENTAL MOVIE 3

Depolymerizing human $\alpha 1A/\beta III$ microtubule imaged by darkfield microscopy at high temporal resolution. Scale bar, 2.5 μm . Recorded at 40 frames per second played back at 30 frames per second.